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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/070,560A

DATE: 10/17/2003
TIME: 14:50:30

Input Set : A:\Sequence.txt
Output Set: N:\CRF4\10172003\J070560A.raw

3 <110> APPLICANT: Entremed, Inc.
4 Liang, Hong
5 Sim, Kim Lee
6 Chang-Murad, Amy
7 Zhou, Xinhua
8 Madsen, John
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10 Bermejo, Lourdes L.
11 Mistry, Firoz Rustom
12 Shepard, Scott R.
13 Schrimsher, Jeffrey L.
15 <120> TITLE OF INVENTION: Method of Producing and Purifying Endostatin Protein
17 <130> FILE REFERENCE: 05213-0551US (43170-258385)
19 <140> CURRENT APPLICATION NUMBER: US 10/070,560A
20 <141> CURRENT FILING DATE: 2003-03-18
22 <150> PRIOR APPLICATION NUMBER: 60/153,698
23 <151> PRIOR FILING DATE: 1999-09-14
25 <150> PRIOR APPLICATION NUMBER: PCT/US00/25166
26 <151> PRIOR FILING DATE: 2000-09-14
28 <160> NUMBER OF SEQ ID NOS: 12
30 <170> SOFTWARE: PatentIn version 3.1
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33 <211> LENGTH: 35
34 <212> TYPE: DNA
35 <213> ORGANISM: Artificial Sequence
37 <220> FEATURE:
38 <223> OTHER INFORMATION: Synthetic primer
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58 <212> TYPE: PRT
59 <213> ORGANISM: Homo sapiens
61 <400> SEQUENCE: 3
63 His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn.

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68	20	25	30
71 Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ala Gly Thr Phe Arg Ala			
72	35	40	45
75 Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala			
76	50	55	60
79 Asp Arg Ala Ala Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe			
80 65	70	75	80
83 Pro Ser Trp Glu Ala Leu Phe Ser Gly Ser Glu Gly Pro Leu Lys Pro			
84	85	90	95
87 Gly Ala Arg Ile Phe Ser Phe Asp Gly Lys Asp Val Leu Arg His Pro			
88	100	105	110
91 Thr Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Asn Gly Arg			
92	115	120	125
95 Arg Leu Thr Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Ala Pro Ser			
96	130	135	140
99 Ala Thr Gly Gln Ala Ser Ser Leu Leu Gly Gly Arg Leu Leu Gly Gln			
100 145	150	155	160
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121 gggctggcgcc gcacccattccg cgccttcctg tcctcgcgc tgcaaggacct gtacagcatc	180		
123 gtgcgcgtg ccgaccgcgc agccgtgccc atcgtcaacc tcaaggacga gctgctgttt	240		
125 cccagctggg aggctctgtt ctcaggctct gagggtccgc tgaagcccg ggcacgcac	300		
127 ttctccctttg acggcaagga cgtcctgagg caccccacct ggccccagaa gagcgtgtgg	360		
129 catggctcgg accccaacgg ggcgcaggctg accgagagct actgtgagac gtggcggacg	420		
131 gaggctccct cggccacggg ccaggcctcc tcgctgctgg gggcaggt cctggggcag	480		
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150	20	25	30
153 Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ala Gly Thr Phe Arg Ala			
154	35	40	45

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157 Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala
158 50 55 60
161 Asp Arg Ala Ala Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe
162 65 70 75 80
165 Pro Ser Trp Glu Ala Leu Phe Ser Gly Ser Glu Gly Pro Leu Lys Pro
166 85 90 95
169 Gly Ala Arg Ile Phe Ser Phe Asp Gly Lys Asp Val Leu Arg His Pro
170 100 105 110
173 Thr Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Asn Gly Arg
174 115 120 125
177 Arg Leu Thr Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Ala Pro Ser
178 130 135 140
181 Ala Thr Gly Gln Ala Ser Ser Leu Leu Gly Gly Arg Leu Leu Gly Gln
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185 Ser Ala Ala Ser Cys His His Ala Tyr Ile Val Leu Cys Ile Glu Asn
186 165 170 175
189 Ser Phe Met Thr Ala Ser
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193 <210> SEQ ID NO: 6
194 <211> LENGTH: 181
195 <212> TYPE: PRT
196 <213> ORGANISM: Homo sapiens
198 <400> SEQUENCE: 6
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205 20 25 30
208 Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ala Gly Thr Phe Arg Ala
209 35 40 45
212 Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala
213 50 55 60
216 Asp Arg Ala Ala Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe
217 65 70 75 80
220 Pro Ser Trp Glu Ala Leu Phe Ser Gly Ser Glu Gly Pro Leu Lys Pro
221 85 90 95
224 Gly Ala Arg Ile Phe Ser Phe Asp Gly Lys Asp Val Leu Arg His Pro
225 100 105 110
228 Thr Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Asn Gly Arg
229 115 120 125
232 Arg Leu Thr Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Ala Pro Ser
233 130 135 140
236 Ala Thr Gly Gln Ala Ser Ser Leu Leu Gly Gly Arg Leu Leu Gly Gln
237 145 150 155 160
240 Ser Ala Ala Ser Cys His His Ala Tyr Ile Val Leu Cys Ile Glu Asn
241 165 170 175
244 Ser Phe Met Thr Ala
245 180
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251 <213> ORGANISM: Homo sapiens
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263 Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ala Gly Thr Phe Arg Ala
264 35 40 45
267 Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala
268 50 55 60
271 Asp Arg Ala Ala Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe
272 65 70 75 80
275 Pro Ser Trp Glu Ala Leu Phe Ser Gly Ser Glu Gly Pro Leu Lys Pro
276 85 90 95
279 Gly Ala Arg Ile Phe Ser Phe Asp Gly Lys Asp Val Leu Arg His Pro
280 100 105 110
283 Thr Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Asn Gly Arg
284 115 120 125
287 Arg Leu Thr Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Ala Pro Ser
288 130 135 140
291 Ala Thr Gly Gln Ala Ser Ser Leu Leu Gly Gly Arg Leu Leu Gly Gln
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295 Ser Ala Ala Ser Cys His His Ala Tyr Ile Val Leu Cys Ile Glu Asn
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304 <211> LENGTH: 179
305 <212> TYPE: PRT
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308 <400> SEQUENCE: 8
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314 Gly Gly Met Arg Gly Ile Arg Gly Ala Asp Phe Gln Cys Phe Gln Gln
315 20 25 30
318 Ala Arg Ala Val Gly Leu Ala Gly Thr Phe Arg Ala Phe Leu Ser Ser
319 35 40 45
322 Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala Asp Arg Ala Ala
323 50 55 60
326 Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe Pro Ser Trp Glu
327 65 70 75 80
330 Ala Leu Phe Ser Gly Ser Glu Gly Pro Leu Lys Pro Gly Ala Arg Ile
331 85 90 95
334 Phe Ser Phe Asp Gly Lys Asp Val Leu Arg His Pro Thr Trp Pro Gln
335 100 105 110
338 Lys Ser Val Trp His Gly Ser Asp Pro Asn Gly Arg Arg Leu Thr Glu
339 115 120 125
342 Ser Tyr Cys Glu Thr Trp Arg Thr Glu Ala Pro Ser Ala Thr Gly Gln

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343	130	135	140	
346	Ala Ser Ser Leu Leu Gly Gly Arg Leu Leu Gly Gln Ser Ala Ala Ser			
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380	accttccgcg ctttcctgtc ctcgcgcctg caggacctgt acagcatcgt gcgcgcgtgcc		180	
382	gaccgcgcag ccgtgcccatt cgtcaacctc aaggacgagc tgctgttcc cagctggag		240	
384	gctctgttct caggctctga gggtccgcgtg aagcccgggg cacgcacatctt ctcccttgac		300	
386	ggcaaggacg tcctgaggca ccccacctgg ccccaagaaga gcgtgtggca tggctcggac		360	
388	cccaacgggc gcaggctgac cgagagctac tgtgagacgt ggcggacgga ggctccctcg		420	
390	gccacgggcc aggccttcctc gctgctgggg ggcaggctcc tggggcagag tgccgcgagc		480	
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407	20	25	30	
410	Ala Arg Ala Val Gly Leu Ala Gly Thr Phe Arg Ala Phe Leu Ser Ser			
411	35	40	45	
414	Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala Asp Arg Ala Ala			
415	50	55	60	
418	Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe Pro Ser Trp Glu			
419	65	70	75	80
422	Ala Leu Phe Ser Gly Ser Glu Gly Pro Leu Lys Pro Gly Ala Arg Ile			
423	85	90	95	
426	Phe Ser Phe Asp Gly Lys Asp Val Leu Arg His Pro Thr Trp Pro Gln			
427	100	105	110	
430	Lys Ser Val Trp His Gly Ser Asp Pro Asn Gly Arg Arg Leu Thr Glu			
431	115	120	125	

VERIFICATION SUMMARYPATENT APPLICATION: **US/10/070,560A**

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